

FIG. 25 is being amended to correct the sequence. Support for this change may be found in (i) the genetic code (see Exhibit A) which indicates the correct amino acids encoded by the codons GAA, GAG, and ACC of FIG. 25 and (ii) the corresponding amino acids in FIG. 11 (see Exhibit B). Note that the truncation 1-163 corresponds to the first 163 amino acids of the mature protein shown in FIG. 11. See page 26, lines 19-21 of the specification. This error was introduced inadvertently and no new matter is being added by the above amendment.

After introducing the above amendments, the Examiner is requested to proceed to examine this case on the merits.

Respectfully submitted,



Robert R. Cook, Ph.D.
Attorney for Applicants
Registration No.: 31,602
Phone No.: 805/447-4955
Date: 2/16/96

Please send future correspondence to:

U. S. Patent Operations/RRC
M/S 10-1-B
AMGEN INC.
Amgen Center
1840 Dehavilland Drive
Thousand Oaks, California 91320-1789

r-HuMGDF (1-163) Translation

ATG	AAA	AGT	CCT	GCA	CCA	CCT	GCA	TGT	GAT	TTA	CGG	GTC	CTG
MET	LYS	SER	PRO	ALA	PRO	PRO	ALA	CYS	ASP	LEU	ARG	VAL	LEU
TCT	AAA	CTG	CTG	CGC	GAC	TCT	CAC	GTG	CTG	CAC	TCT	CGT	CTG
SER	LYS	LEU	LEU	ARG	ASP	SER	HIS	VAL	LEU	HIS	SER	ARG	LEU
TCC	CAG	TGC	CCG	GAA	GTT	CAC	CCG	CTG	CCG	ACC	CCG	GTT	CTG
SER	GLN	CYS	PRO	GLU	VAL	HIS	PRO	LEU	PRO	THR	PRO	VAL	LEU
CTT	CCG	GCT	GTC	GAC	TTC	TCC	CTG	GGT	GAA	TGG	AAA	ACC	CAG
LEU	PRO	ALA	VAL	ASP	PHE	SER	LEU	GLY	GLU	TRP	LYS	THR	GLN
ATG	GAA	GAG	ACC	AAA	GCT	CAG	GAC	ATC	CTG	GGT	GCA	GTA	ACT
MET	ALA	ALA	ARG	LYS	ALA	GLN	ASP	ILE	LEU	GLY	ALA	VAL	THR
	GLU	GLU	THR										
CTG	CTT	CTG	GAA	GGC	GTT	ATG	GCT	GCA	CGT	GGC	CAG	CTT	GGC
LEU	LEU	LEU	GLU	GLY	VAL	MET	ALA	ALA	ARG	GLY	GLN	LEU	GLY
CCG	ACC	TGC	CTG	TCT	TCC	CTG	CTT	GGC	CAG	CTG	TCT	GGC	CAG
PRO	THR	CYS	LEU	SER	SER	LEU	LEU	GLY	GLN	LEU	SER	GLY	GLN
GTT	CGT	CTG	CTG	CTC	GGC	GCT	CTG	CAG	TCT	CTG	CTT	GGC	ACC
VAL	ARG	LEU	LEU	LEU	GLY	ALA	LEU	GLN	SER	LEU	LEU	GLY	THR
CAG	CTG	CCG	CCA	CAG	GGC	CGT	ACC	ACT	GCT	CAC	AAG	GAT	CCG
GLN	LEU	PRO	PRO	GLN	GLY	ARG	THR	THR	ALA	HIS	LYS	ASP	PRO
AAC	GCT	ATC	TTC	CTG	TCT	TTC	CAG	CAC	CTG	CTG	CGT	GGC	AAA
ASN	ALA	ILE	PHE	LEU	SER	PHE	GLN	HIS	LEU	LEU	ARG	GLY	LYS
GTT	CGT	TTC	CTG	ATG	CTG	GTT	GGC	GGT	TCT	ACC	CTG	TGC	GTT
VAL	ARG	PHE	LEU	MET	LEU	VAL	GLY	GLY	SER	THR	LEU	CYS	VAL
CGT	CGG	GCG	CCG	CCA	ACC	ACT	GCT	GTT	CCG	TCT	TAA		
ARG	ARG	ALA	PRO	PRO	THR	THR	ALA	VAL	PRO	SER	STOP		

Figure 25